SEQUENCE LISTING

<110> KISHIMOTO, Takahide
SOGABE, Atsushi
OKA, Masanori

<120> MODIFIED SARCOSINE OXIDASE, PROCESS FOR PRODUCING THE SAME AND REAGENT COMPOSITION USING THE SAME

<130> 235670

<150> PCT/JP2003/014423
<151> 2003-11-13

<150> JP2002-329427
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<150> JP2002-329428
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<151> 2003-02-12

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<170> PatentIn version 3.1

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35 40 45

Gly Asp Thr Arg Ile Ile Arg His Ala Tyr Gly Glu Gly Arg Glu Tyr
50 55 60

Val Pro Phe Ala Leu Arg Ala Gln Glu Leu Trp Tyr Glu Leu Glu Lys
65 70 75 80

Glu Thr His His Lys Ile Phe Thr Lys Thr Gly Val Leu Val Phe Gly
85 90 95

Pro Lys Gly Glu Ala Pro Phe Val Ala Glu Thr Met Glu Ala Ala Lys
100 105 110

Glu His Ser Leu Asp Val Asp Leu Leu Glu Gly Ser Glu Ile Asn Lys
115 120 125

Arg Trp Pro Gly Val Thr Val Pro Glu Asn Tyr Asn Ala Ile Phe Glu
130 135 140

Lys Asn Ser Gly Val Leu Phe Ser Glu Asn Cys Ile Arg Ala Tyr Arg
145 150 155 160

Glu Leu Ala Glu Ala Asn Gly Ala Lys Val Leu Thr Tyr Thr Pro Val
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Gly Ser Phe Thr Ala Ser Lys Leu Ile Val Ser Met Gly Ala Trp Asn
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Ser Lys Leu Leu Ser Lys Leu Asn Ile Glu Ile Pro Leu Gln Pro Tyr
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Arg Gln Val Val Gly Phe Phe Glu Cys Asp Glu Lys Lys Tyr Ser Asn
225 230 235 240

Thr His Gly Tyr Pro Ala Phe Met Val Glu Val Pro Thr Gly Ile Tyr
245 250 255

Tyr Gly Phe Pro Ser Phe Gly Gly Cys Gly Leu Lys Ile Gly Tyr His
260 265 270

Thr Tyr Gly Gln Lys Ile Asp Pro Asp Thr Ile Asn Arg Glu Phe Gly
275 280 285

Ile Tyr Pro Glu Asp Glu Gly Asn Ile Arg Lys Phe Leu Glu Thr Tyr
290 295 300

Met Pro Gly Ala Thr Gly Glu Leu Lys Ser Gly Ala Val Cys Met Tyr
305 310 315 320

Thr Lys Thr Pro Asp Glu His Phe Val Ile Asp Leu His Pro Gln Phe
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Met Gly Met Ala Ala Gly Tyr Tyr Leu Ser Lys Gln Gly Val Lys Thr
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cta ttg gta gat tca ttt cat cct ccc cat aca aat ggc agc cat cat      144
Leu Leu Val Asp Ser Phe His Pro Pro His Thr Asn Gly Ser His His
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ggc gat aca cgg atc att cgt cac gca tat ggc gaa gga aga gag tat      192
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gta ccg ttt gcc ttg aga gca caa gag tta tgg tat gaa tta gaa aag      240
Val Pro Phe Ala Leu Arg Ala Gln Glu Leu Trp Tyr Glu Leu Glu Lys
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gag act cat cat aaa ata ttt aca aaa aca ggt gta ctc gtt ttt ggt      288
Glu Thr His His Lys Ile Phe Thr Lys Thr Gly Val Leu Val Phe Gly
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cct aaa gga gaa gct cct ttc gtt gcc gaa aca atg gaa gcc gca aag      336
Pro Lys Gly Glu Ala Pro Phe Val Ala Glu Thr Met Glu Ala Ala Lys
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Glu His Ser Leu Asp Val Asp Leu Leu Glu Gly Ser Glu Ile Asn Lys
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Arg Trp Pro Gly Val Thr Val Pro Glu Asn Tyr Asn Ala Ile Phe Glu
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aaa aat tct ggt gtc tta ttt agt gaa aat tgt att cgc gct tac cgt      480
Lys Asn Ser Gly Val Leu Phe Ser Glu Asn Cys Ile Arg Ala Tyr Arg
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gaa ttg gcg gaa gca aat ggt gcg aaa gtt cta acg tac aca ccc gtt      528
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Gly Ser Phe Thr Ala Ser Lys Leu Ile Val Ser Met Gly Ala Trp Asn
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agc aaa ctg cta tca aaa tta aat att gaa atc cca ttg cag cca tac      672
Ser Lys Leu Leu Ser Lys Leu Asn Ile Glu Ile Pro Leu Gln Pro Tyr
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aca cat ggt tat ccg gcg ttc atg gtc gaa gtc cca act ggc atc tat      768
Thr His Gly Tyr Pro Ala Phe Met Val Glu Val Pro Thr Gly Ile Tyr
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Tyr Gly Phe Pro Ser Phe Gly Gly Cys Gly Leu Lys Ile Gly Tyr His
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acg tat ggt caa aaa atc gat cca gat acg att aat cgt gaa ttt ggt      864
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att tac ccg gag gat gaa ggg aat att cgc aaa ttc ctg gaa aca tat      912
Ile Tyr Pro Glu Asp Glu Gly Asn Ile Arg Lys Phe Leu Glu Thr Tyr
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aca aaa aca cct gat gag cat ttc gtg att gat tta cat cct caa ttc     1008
Thr Lys Thr Pro Asp Glu His Phe Val Ile Asp Leu His Pro Gln Phe
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Ser Asn Val Ala Ile Ala Ala Gly Phe Ser Gly His Gly Phe Lys Phe
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Ser Ser Val Val Gly Glu Thr Leu Ser Gln Leu Ala Val Thr Gly Lys
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Thr Glu His Asp Ile Ser Ile Phe Ser Ile Asn Arg Pro Ala Leu Lys
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caa aaa gaa acg att                                                1167
Gln Lys Glu Thr Ile
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